36



SEQUENCE LISTING

<110> Xia, Yu-Ping et al.

<120> METHODS FOR TREATING INFLAMMATORY SKIN DISEASES

<130> REG 710b

<140> 09/773,877

<141> 2001-01-31

<160> 27

<170> PatentIn version 3.0

<210> 1

<211> 36

<212> DNA

<213> Artificial

<220>

<223> Primer

<400> 1

gactagcagt ccggaggtag acctttcgta gagatg

<210> 2

<211> 33

<212> DNA

<213> Artificial

<220>		
<223>	Primer	
<400> cggact	2 caga accacatcta tgattgtatt ggt	33
<210>	3	
<211>	35	
<212>	DNA	
<213>	Artificial	
<220>		
<223>	Primer	
<400>		35
acaacc	atag atgtggttct gagtccgtct catgg	,,
<210>	4	
<211>	38	
<212>	DNA	
<213>	Artificial	
<220>		
<223>	Primer	
<400>		38
gacaac	gece gggeeetttt catggaeeet gacaaatg	30
<210>	5	
<211>	36	
<212>	DNA	
<213>	Artificial	
<220>		

<223> Primer

gactag	cagt ccggaggtag acct	ttcgta gagatg	36
<210>	6		
<211>	38		
<212>	DNA		
<213>	Artificial		
<220>			
<223>	Primer		
<400>	6 ggca acagctggat atcta	atgatt gtattggt	38
ccccg	ggod dodgooggdo dooo	acgace geacegge	
<210>	7		
<211>	39		
<212>	DNA		
<213>	Artificial		•
<220>			
<223>	Primer		
<400>	7 ctgt tgcccaggaa gtcgo	ctggag ctgctggta	39
<210>	8		
<211>	39		
<212>	DNA		
<213>	Artificial		
<220>			
<223>	Primer		
<400> attttca	8 atgc acaatgacct cggts	getete eegaaateg	39

```
<210> 9
 <211> 38
 <212> DNA
 <213> Artificial
 <220>
  <223> Primer
 <400> 9
 tcatagatat ccagctgttg cccaggaagt cgctggag
 <210> 10
 <211> 39
 <212> DNA
 <213> Artificial
 <220>
 <223> Primer
 <400> 10
 gataatgccc gggccatttt catgcacaat gacctcggt
 <210> 11
 <211> 1704
 <212> DNA
<213> Artificial
 <220>
```

<223> Flt1(1-3)-Fc

<222> (1)..(1704)

<220>

<221> CDS

4

38

39

<400>	11														
atg gt Met Va 1	c agc l Ser	tac Tyr	tgg Trp 5	gac Asp	acc Thr	ggg Gly	gtc Val	ctg Leu 10	ctg Leu	Cys	gcg Ala	Leu	Leu 15	agc Ser	48
tgt ct Cys Le	g ctt u Leu	ctc Leu 20	aca Thr	gga Gly	tct Ser	agt Ser	tca Ser 25	ggt Gly	tca Ser	aaa Lys	tta Leu	aaa Lys 30	gat Asp	cct Pro	96
gaa ct Glu Le	g agt u Ser 35	tta Leu	aaa Lys	ggc Gly	acc Thr	cag Gln 40	cac His	atc Ile	atg Met	caa Gln	gca Ala 45	ggc	cag Gln	aca Thr	144
ctg ca Leu Hi 50	t ctc s Leu	caa Gln	tgc Cys	agg Arg	999 Gly 55	gaa Glu	gca Ala	gcc Ala	cat His	aaa Lys 60	tgg Trp	tct Ser	ttg Leu	cct Pro	192
gaa at Glu Me 65															240
tgt gg Cys Gl															288
gct ca Ala Gl															336
cct ac Pro Th		Lys													384
agt ga Ser As 13	p Thr														432
att at Ile Il 145		_		_					_			_		_	480
acg tc Thr Se															528
ttg at Leu Il															576
atc at Ile Il															624
gca ac Ala Th 21	r Val														672
caa ac	c aat	aca	atc	ata	gat	gtc	caa	ata	agc	aca	cca	cgc	cca	gtc	720

Gln 225	Thr	Asn	Thr	Ile	Ile 230	Asp	Val	Gln	Ile	Ser 235	Thr	Pro	Arg	Pro	Val 240	
						act Thr										768
						caa Gln										816
						agg Arg										864
						gtt Val 295										912
						tgt Cys										960
						cat His										1008
			_	_		act Thr			_		_	_		_		1056
						tca Ser										1104
gac Asp	acc Thr 370	ctc Leu	atg Met	atc Ile	tcc Ser	cgg Arg 375	acc Thr	cct Pro	gag Glu	gtc Val	aca Thr 380	tgc Cys	gtg Val	gtg Val	gtg Val	1152
						cct Pro										1200
						gcc Ala										1248
aac Asn	agc Ser	acg Thr	tac Tyr 420	cgt Arg	gtg Val	gtc Val	agc Ser	gtc Val 425	ctc Leu	acc Thr	gtc Val	ctg Leu	cac His 430	cag Gln	gac Asp	1296
						tac Tyr										1344
cca	gcc	ccc	atc	gag	aaa	acc	atc	tcc	aaa	gcc	aaa	ggg	cag	ccc	cga	1392

Pro Ala Pro I 450	le Glu Lys	Thr Ile Se	er Lys Ala	Lys Gly Gln 460	Pro Arg	
gaa cca cag g Glu Pro Gln V 465						1440
aac cag gtc a Asn Gln Val S	gc ctg acc er Leu Thr 485	tgc ctg gt Cys Leu Va	tc aaa ggc al Lys Gly 490	ttc tat ccc Phe Tyr Pro	agc gac Ser Asp 495	1488
atc gcc gtg g Ile Ala Val G 5		Ser Asn G				1536
acc acg cct c Thr Thr Pro P 515						1584
aag ctc acc g Lys Leu Thr V 530						1632
tgc tcc gtg a Cys Ser Val M 545						1680
ctc tcc ctg to Leu Ser Leu So		_				1704
<210> 12						
<211> 567						
<212> PRT						
<213> Artific	cial					
<400> 12						
Met Val Ser Ty 1	yr Trp Asp 5	Thr Gly Va	al Leu Leu 10	Cys Ala Leu	Leu Ser 15	
Cys Leu Leu Le 20		Ser Ser Se 25		Lys Leu Lys 30	Asp Pro	
Glu Leu Ser Le 35	eu Lys Gly	Thr Gln Hi	is Ile Met	Gln Ala Gly 45	Gln Thr	

Leu His Leu Gln Cys Arg Gly Glu Ala Ala His Lys Trp Ser Leu Pro 55 Glu Met Val Ser Lys Glu Ser Glu Arg Leu Ser Ile Thr Lys Ser Ala Cys Gly Arg Asn Gly Lys Gln Phe Cys Ser Thr Leu Thr Leu Asn Thr Ala Gln Ala Asn His Thr Gly Phe Tyr Ser Cys Lys Tyr Leu Ala Val 100 105 Pro Thr Ser Lys Lys Glu Thr Glu Ser Ala Ile Tyr Ile Phe Ile 115 Ser Asp Thr Gly Arg Pro Phe Val Glu Met Tyr Ser Glu Ile Pro Glu 130 135 Ile Ile His Met Thr Glu Gly Arg Glu Leu Val Ile Pro Cys Arg Val Thr Ser Pro Asn Ile Thr Val Thr Leu Lys Lys Phe Pro Leu Asp Thr 170 165 Leu Ile Pro Asp Gly Lys Arg Ile Ile Trp Asp Ser Arg Lys Gly Phe 185 Ile Ile Ser Asn Ala Thr Tyr Lys Glu Ile Gly Leu Leu Thr Cys Glu 195 200 Ala Thr Val Asn Gly His Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg 210 215 Gln Thr Asn Thr Ile Ile Asp Val Gln Ile Ser Thr Pro Arg Pro Val Lys Leu Leu Arg Gly His Thr Leu Val Leu Asn Cys Thr Ala Thr Thr 245 Pro Leu Asn Thr Arg Val Gln Met Thr Trp Ser Tyr Pro Asp Glu Lys 260 265

Asn Lys Arg Ala Ser Val Arg Arg Ile Asp Gln Ser Asn Ser His

275 280 285

АІА	290	iie	Pile	IYL	Ser	295	Бец	1111	116	Asp	300	мес	GIII	ASII	цуг
Asp 305	Lys	Gly	Leu	Tyr	Thr 310	Cys	Arg	Val	Arg	Ser 315	Gly	Pro	Ser	Phe	Lys 320
Ser	Val	Asn	Thr	Ser 325	Val	His	Ile	Tyr	Asp 330	Lys	Ala	Gly	Pro	Gly 335	Gli
Pro	Lys	Ser	Cys 340	Asp	Lys	Thr	His	Thr 345	Cys	Pro	Pro	Cys	Pro 350	Ala	Pro
Glu	Leu	Leu 355	Gly	Gly	Pro	Ser	Val 360	Phe	Leu	Phe	Pro	Pro 365	Lys	Pro	Lys
Asp	Thr 370	Leu	Met	Ile	Ser	Arg 375	Thr	Pro	Glu	Val	Thr 380	Cys	Val	Val	Va]
Asp 385	Val	Ser	His	Glu	Asp 390	Pro	Glu	Val	Lys	Phe 395	Asn	Trp	Tyr	Val	Asp 400
Gly	Val	Glu	Val	His 405	Asn	Ala	Lys	Thr	Lys 410	Pro	Arg	Glu	Glu	Gln 415	Туг
Asn	Ser	Thr	Tyr 420	Arg	Val	Val	Ser	Val 425	Leu	Thr	Val	Leu	His 430	Gln	Asp
Trp	Leu	Asn 435	Gly	Lys	Glu	Tyr	Lys 440	Cys	Lys	Val	Ser	Asn 445	Lys	Ala	Leu
Pro	Ala 450	Pro	Ile	Glu	Lys	Thr 455	Ile	Ser	Lys	Ala	Lys 460	Gly	Gln	Pro	Arg
Glu 465	Pro	Gln	Val	Tyr	Thr 470	Leu	Pro	Pro	Ser	Arg 475	Asp	Glu	Leu	Thr	Lys 480
Asn	Gln	Val	Ser	Leu 485	Thr	Cys	Leu	Val	Lys 490	Gly	Phe	Tyr	Pro	Ser 495	Asp
Ile	Ala	Val	Glu 500	Trp	Glu	Ser	Asn	Gly 505	Gln	Pro	Glu	Asn	Asn 510	Tyr	Lys

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser 520 Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser 530 Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser 555 545 550 Leu Ser Leu Ser Pro Gly Lys <210> 13 <211> 1674 <212> DNA <213> Artificial <220> <223> Mut 1 <220> <221> CDS <222> (1)..(1674) <400> 13 atg gtc agc tac tgg gac acc ggg gtc ctg ctg tgc gcg ctg ctc agc 48 Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser tgt ctg ctt ctc aca gga tct agt tca ggt tca aaa tta aaa gat cct 96 Cys Leu Leu Thr Gly Ser Ser Gly Ser Lys Leu Lys Asp Pro gaa ctg agt tta aaa ggc acc cag cac atc atg caa gca ggc cag aca 144 Glu Leu Ser Leu Lys Gly Thr Gln His Ile Met Gln Ala Gly Gln Thr ctg cat ctc caa tgc agg ggg gaa gca gcc cat aaa tgg tct ttg cct 192 Leu His Leu Gln Cys Arg Gly Glu Ala Ala His Lys Trp Ser Leu Pro 55 gaa atg gtg agt aag gaa agc gaa agg ctg agc ata act aaa tct gcc 240 Glu Met Val Ser Lys Glu Ser Glu Arg Leu Ser Ile Thr Lys Ser Ala

75

70

						caa Gln										288
						ggc Gly										336
cct Pro	act Thr	tca Ser 115	aag Lys	aag Lys	aag Lys	gaa Glu	aca Thr 120	gaa Glu	tct Ser	gca Ala	atc Ile	tat Tyr 125	ata Ile	ttt Phe	att Ile	384
						ttc Phe 135										432
att Ile 145	ata Ile	cac His	atg Met	act Thr	gaa Glu 150	gga Gly	agg Arg	gag Glu	ctc Leu	gtc Val 155	att Ile	ccc Pro	tgc Cys	cgg Arg	gtt Val 160	480
						gtt Val										528
ttg Leu	atc Ile	cct Pro	gat Asp 180	gga Gly	aaa Lys	cgc Arg	ata Ile	atc Ile 185	tgg Trp	gac Asp	agt Ser	aga Arg	aag Lys 190	ggc	ttc Phe	576
						tac Tyr										624
						ttg Leu 215										672
						gat Asp										720
						act Thr										768
			_	_	_	caa Gln	_			_			_	_		816
						gcc Ala										864
						gac Asp 295										912
agt Ser	gga Gly	cca Pro	tca Ser	ttc Phe	aaa Lys	tct Ser	gtt Val	aac Asn	acc Thr	tca Ser	gtg Val	cat His	ata Ile	tat Tyr	gat Asp	960

305					310					315					320		
											aaa Lys					10	80
											ccg Pro					10	56
ttc Phe	ccc Pro	cca Pro 355	aaa Lys	ccc Pro	aag Lys	gac Asp	acc Thr 360	ctc Leu	atg Met	atc Ile	tcc Ser	cgg Arg 365	acc Thr	cct Pro	gag Glu	11	04
gtc Val	aca Thr 370	tgc Cys	gtg Val	gtg Val	gtg Val	gac Asp 375	gtg Val	agc Ser	cac His	gaa Glu	gac Asp 380	cct Pro	gag Glu	gtc Val	aag Lys	11	52
											aat Asn					12	00
											gtg Val					12	48
acc Thr	gtc Val	ctg Leu	cac His 420	cag Gln	gac Asp	tgg Trp	ctg Leu	aat Asn 425	ggc Gly	aag Lys	gag Glu	tac Tyr	aag Lys 430	tgc Cys	aag Lys	12	96
_				_							aaa Lys					13	44
											acc Thr 460					13	92
											acc Thr					14	40
				_	_		_				gag Glu	_			_	14	88
_					_		_				ctg Leu	_		_		15	36
					_	_				_	aag Lys	_				15	84
											gag Glu 540					16	32

1674

cac tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa tga His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys 545 550 555

<210> 14

<211> 557

<212> PRT

<213> Artificial

<400> 14

Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser 1 5 10 15

Cys Leu Leu Thr Gly Ser Ser Ser Gly Ser Lys Leu Lys Asp Pro 20 25 30

Glu Leu Ser Leu Lys Gly Thr Gln His Ile Met Gln Ala Gly Gln Thr 35 40 45

Leu His Leu Gln Cys Arg Gly Glu Ala Ala His Lys Trp Ser Leu Pro 50 55 60

Glu Met Val Ser Lys Glu Ser Glu Arg Leu Ser Ile Thr Lys Ser Ala 65 70 75 80

Cys Gly Arg Asn Gly Lys Gln Phe Cys Ser Thr Leu Thr Leu Asn Thr 85 90 95

Ala Gln Ala Asn His Thr Gly Phe Tyr Ser Cys Lys Tyr Leu Ala Val 100 105 110

Pro Thr Ser Lys Lys Glu Thr Glu Ser Ala Ile Tyr Ile Phe Ile 115 120 125

Ser Asp Thr Gly Arg Pro Phe Val Glu Met Tyr Ser Glu Ile Pro Glu 130 135 140

Ile Ile His Met Thr Glu Gly Arg Glu Leu Val Ile Pro Cys Arg Val
145 150 155 160

Thr	Ser	Pro	Asn	Ile 165	Thr	Val	Thr	Leu	Lys 170	Lys	Phe	Pro	Leu	Asp 175	Thr
Leu	Ile	Pro	Asp 180	Gly	Lys	Arg	Ile	Ile 185	Trp	Asp	Ser	Arg	Lys 190	Gly	Phe
Ile	Ile	Ser 195	Asn	Ala	Thr	Tyr	Lys 200	Glu	Ile	Gly	Leu	Leu 205	Thr	Cys	Glu
Ala	Thr 210	Val	Asn	Gly	His	Leu 215	Tyr	Lys	Thr	Asn	Tyr 220	Leu	Thr	His	Arg
Gln 225	Thr	Asn	Thr	Ile	Ile 230	Asp	Val	Gln	Ile	Ser 235	Thr	Pro	Arg	Pro	Val 240
Lys	Leu	Leu	Arg	Gly 245	His	Thr	Leu	Val	Leu 250	Asn	Cys	Thr	Ala	Thr 255	Thr
Pro	Leu	Asn	Thr 260	Arg	Val	Gln	Met	Thr 265	Trp	Ser	Tyr	Pro	Asp 270	Glu	Ile
Asp	Gln	Ser 275	Asn	Ser	His	Ala	Asn 280	Ile	Phe	Tyr	Ser	Val 285	Leu	Thr	Ile
Asp	Lys 290	Met	Gln	Asn	Lys	Asp 295	Lys	Gly	Leu	Tyr	Thr 300	Cys	Arg	Val	Arg
Ser 305	Gly	Pro	Ser	Phe	Lys 310	Ser	Val	Asn	Thr	Ser 315	Val	His	Ile	Tyr	Asp 320
Lys	Ala	Gly	Pro	Gly 325	Glu	Pro	Lys	Ser	Cys 330	Asp	Lys	Thr	His	Thr 335	Cys
Pro	Pro	Cys	Pro 340	Ala	Pro	Glu	Leu	Leu 345	Gly	Gly	Pro	Ser	Val 350	Phe	Leu
Phe	Pro	Pro 355	Lys	Pro	Lys	Asp	Thr 360	Leu	Met	Ile	Ser	Arg 365	Thr	Pro	Glu
Val	Thr 370	Cys	Val	Val	Val	Asp 375	Val	Ser	His	Glu	Asp 380	Pro	Glu	Val	Lys
Phe 385	Asn	Trp	Tyr	Val	Asp 390	Gly	Val	Glu	Val	His 395	Asn	Ala	Lys	Thr	Lys 400

Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu 405 410 415

Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys 420 425 430

Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys 435 440 445

Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser 450 455 460

Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys 465 470 475 480

Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln 485 490 495

Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly 500 505 510

Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln 515 520 525

Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn 530 540

His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys 545 550 555

<210> 15

<211> 1359

<212> DNA

<213> Artificial

<220>

<223> Mut2

<220>

<221> CDS

<40	0 > :	15															
														ctc Leu 15			48
														gta Val			96
														agg Arg		1	44
														act Thr		1	92
														ata Ile		2	40
		_	_	_							_	_		aaa Lys 95	-	`2	88
														tat Tyr		3	36
						_							_	gtc Val		3	84
														ctt Leu		4	32
		_		_				_		_	_	_		atg Met		4	80
														aac Asn 175		5	28
		_	_				_		_	_			_	aaa Lys		5	76
														gtt Val		6	24
acc	tca	gtg	cat	ata	tat	gat	aaa	gca	ggc	ccg	ggc	gag	ccc	aaa	tct	6	72

Thr	Ser 210	Val	His	Ile	Tyr	Asp 215	Lys	Ala	Gly	Pro	Gly 220	Glu	Pro	Lys	Ser	
								ccg Pro								720
								ccc Pro								768
atg Met	atc Ile	tcc Ser	cgg Arg 260	acc Thr	cct Pro	gag Glu	gtc Val	aca Thr 265	tgc Cys	gtg Val	gtg Val	gtg Val	gac Asp 270	gtg Val	agc Ser	816
								aac Asn								864
								cgg Arg								912
								gtc Val								960
								tcc Ser								1008
								aaa Lys 345								1056
								gat Asp								1104
								ttc Phe								1152
			_			_	_	gag Glu				_		_		1200
								ttc Phe								1248
								999 Gly 425								1296
_			_	_				tac Tyr	_	_	_	_				1344

tct ccg ggt aaa tga Ser Pro Gly Lys 450

<210> 16

<211> 452

<212> PRT

<213> Artificial

<400> 16

Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser 1 5 10 15

Cys Leu Leu Thr Gly Ser Ser Ser Gly Gly Arg Pro Phe Val Glu 20 25 30

Met Tyr Ser Glu Ile Pro Glu Ile Ile His Met Thr Glu Gly Arg Glu 35 40 45

Leu Val Ile Pro Cys Arg Val Thr Ser Pro Asn Ile Thr Val Thr Leu 50 55 60

Lys Lys Phe Pro Leu Asp Thr Leu Ile Pro Asp Gly Lys Arg Ile Ile 65 70 75 80

Trp Asp Ser Arg Lys Gly Phe Ile Ile Ser Asn Ala Thr Tyr Lys Glu 85 90 95

Ile Gly Leu Leu Thr Cys Glu Ala Thr Val Asn Gly His Leu Tyr Lys
100 105 110

Thr Asn Tyr Leu Thr His Arg Gln Thr Asn Thr Ile Ile Asp Val Gln
115 120 125

Ile Ser Thr Pro Arg Pro Val Lys Leu Leu Arg Gly His Thr Leu Val 130 135 140

Leu Asn Cys Thr Ala Thr Thr Pro Leu Asn Thr Arg Val Gln Met Thr 145 150 155 160

Trp Ser Tyr Pro Asp Glu Ile Asp Gln Ser Asn Ser His Ala Asn Ile

Phe	Tyr	Ser	Val	Leu	Thr	Ile	Asp	Lys	Met	Gln	Asn	Lys	Asp	Lys	Gly
			180					185					190		

- Leu Tyr Thr Cys Arg Val Arg Ser Gly Pro Ser Phe Lys Ser Val Asn 195 200 205
- Thr Ser Val His Ile Tyr Asp Lys Ala Gly Pro Gly Glu Pro Lys Ser 210 215 220
- Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu 225 230 235 240
- Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu 245 250 255
- Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser 260 265 270
- His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu 275 280 285
- Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr 290 295 300
- Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn 305 310 315 320
- Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro 325 330 335
- Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln 340 345 350
- Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val 355 360 365
- Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val 370 375 380
- Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro 385 390 395 400

Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr 405 410 Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val 425 Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu 440 Ser Pro Gly Lys 450 <210> 17 <211> 1389 <212> DNA <213> Artificial <220> <223> Mut3 <220> <221> CDS <222> (1)..(1389) <400> 17 atg gtc agc tac tgg gac acc ggg gtc ctg ctg tgc gcg ctg ctc agc 48 Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser tgt ctg ctt ctc aca gga tct agt tcc gga ggt aga cct ttc gta gag 96 Cys Leu Leu Thr Gly Ser Ser Gly Gly Arg Pro Phe Val Glu atg tac agt gaa atc ccc gaa att ata cac atg act gaa gga agg gag 144 Met Tyr Ser Glu Ile Pro Glu Ile Ile His Met Thr Glu Gly Arg Glu ctc gtc att ccc tgc cgg gtt acg tca cct aac atc act gtt act tta 192 Leu Val Ile Pro Cys Arg Val Thr Ser Pro Asn Ile Thr Val Thr Leu 55 aaa aag ttt cca ctt gac act ttg atc cct gat gga aaa cgc ata atc 240 Lys Lys Phe Pro Leu Asp Thr Leu Ile Pro Asp Gly Lys Arg Ile Ile

75

70

		ggc Gly								288
		tgt Cys								336
		cat His								384
		cca Pro								432
		acc Thr 150								480
		gaa Glu								528
		tcc Ser								576
		aac Asn								624
 _		ttc Phe			_					672
		ggc Gly 230								720
		gca Ala								768
		ccc Pro	_	_		_				816
		gtg Val								864
		gtg Val								912

aag Lys 305	ccg Pro	cgg Arg	gag Glu	gag Glu	cag Gln 310	tac Tyr	aac Asn	agc Ser	acg Thr	tac Tyr 315	cgt Arg	gtg Val	gtc Val	agc Ser	gtc Val 320	960
					cag Gln											1008
aag Lys	gtc Val	tcc Ser	aac Asn 340	aaa Lys	gcc Ala	ctc Leu	cca Pro	gcc Ala 345	ccc Pro	atc Ile	gag Glu	aaa Lys	acc Thr 350	atc Ile	tcc Ser	1056
aaa Lys	gcc Ala	aaa Lys 355	gly aaa	cag Gln	ccc Pro	cga Arg	gaa Glu 360	cca Pro	cag Gln	gtg Val	tac Tyr	acc Thr 365	ctg Leu	ccc Pro	cca Pro	1104
tcc Ser	cgg Arg 370	gat Asp	gag Glu	ctg Leu	acc Thr	aag Lys 375	aac Asn	cag Gln	gtc Val	agc Ser	ctg Leu 380	acc Thr	tgc Cys	ctg Leu	gtc Val	1152
					agc Ser 390											1200
cag Gln	ccg Pro	gag Glu	aac Asn	aac Asn 405	tac Tyr	aag Lys	acc Thr	acg Thr	cct Pro 410	ccc Pro	gtg Val	ctg Leu	gac Asp	tcc Ser 415	gac Asp	1248
					tac Tyr											1296
					ttc Phe											1344
					aag Lys									tga		1389

<210> 18

<211> 462

<212> PRT

<213> Artificial

<400> 18

Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser 1 5 10 15

Cys Leu Leu Thr Gly Ser Ser Ser Gly Gly Arg Pro Phe Val Glu 25 Met Tyr Ser Glu Ile Pro Glu Ile Ile His Met Thr Glu Gly Arg Glu 40 Leu Val Ile Pro Cys Arg Val Thr Ser Pro Asn Ile Thr Val Thr Leu 50 55 Lys Lys Phe Pro Leu Asp Thr Leu Ile Pro Asp Gly Lys Arg Ile Ile 70 75 Trp Asp Ser Arg Lys Gly Phe Ile Ile Ser Asn Ala Thr Tyr Lys Glu 85 Ile Gly Leu Leu Thr Cys Glu Ala Thr Val Asn Gly His Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg Gln Thr Asn Thr Ile Ile Asp Val Gln 120 Ile Ser Thr Pro Arg Pro Val Lys Leu Leu Arg Gly His Thr Leu Val 130 135 Leu Asn Cys Thr Ala Thr Thr Pro Leu Asn Thr Arg Val Gln Met Thr 150 155 Trp Ser Tyr Pro Asp Glu Lys Asn Lys Arg Ala Ser Val Arg Arg Arg 165 170 Ile Asp Gln Ser Asn Ser His Ala Asn Ile Phe Tyr Ser Val Leu Thr Ile Asp Lys Met Gln Asn Lys Asp Lys Gly Leu Tyr Thr Cys Arg Val 195 Arg Ser Gly Pro Ser Phe Lys Ser Val Asn Thr Ser Val His Ile Tyr 210 215 Asp Lys Ala Gly Pro Gly Glu Pro Lys Ser Cys Asp Lys Thr His Thr 240 225 230 235

Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe

Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro 260 265 270

Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val 275 280 285

Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr 290 295 300

Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val 305 310 315 320

Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys 325 330 335

Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser 340 345 350

Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro 355 360 365

Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val 370 380

Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly 385 390 395 400

Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp 405 410 415

Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp 420 425 430

Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His 435 440 445

Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys 450 455 460

<210> 19

<211> 1704

<212> DNA

<213> Artificial

<220>

<223> Mut 4

<220>

<221> CDS

<222> (1)..(1704)

<400)> :	19														
					gac Asp											48
_	_				gga Gly		_									96
					ggc Gly											144
_					agg Arg											192
_	_		_	_	gaa Glu 70	_	_		_	_					_	240
_		_			aaa Lys			_	_				_			288
					act Thr				_	_						336
			_	_	aag Lys	_		_		_						384
_	_			_	cct Pro		_		_		_				_	432
att	ata	cac	atg	act	gaa	gga	agg	gag	ctc	gtc	att	CCC	tgc	cgg	gtt	480

Ile 145	Ile	His	Met	Thr	Glu 150	Gly	Arg	Glu	Leu	Val 155	Ile	Pro	Cys	Arg	Val 160	
						gtt Val										528
						cgc Arg										576
						tac Tyr										624
						ttg Leu 215										672
						gat Asp										720
						act Thr										768
	-		_	_	_	caa Gln	_									816
						agg Arg										864
						gtt Val 295										912
_						tgt Cys	_	_								960
	_					cat His			_		_		_		-	1008
			_	_		act Thr			_		_	_		_		1056
						tca Ser										1104
gac	acc	ctc	atg	atc	tcc	cgg	acc	cct	gag	gtc	aca	tgc	gtg	gtg	gtg	1152

Asp	Thr 370	Leu	Met	Ile	Ser	Arg 375	Thr	Pro	Glu	Val	Thr 380	Cys	Val	Val	Val		
								gtc Val								<u>.</u>	1200
								aca Thr								=	1248
								gtc Val 425								:	1296
								tgc Cys								=	1344
								tcc Ser								=	1392
								cca Pro								-	1440
	_	_	_	_				gtc Val								:	1488
								999 Gly 505								1	1536
	_				_	_		gac Asp							_	1	1584
								tgg Trp								1	1632
								cac His								1	1680
					ggt Gly		tga									1	1704

<210> 20

<211> 567

<212> PRT

<213> Artificial

<400> 20

Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser 1 5 10 15

Cys Leu Leu Thr Gly Ser Ser Ser Gly Ser Lys Leu Lys Asp Pro 20 25 30

Glu Leu Ser Leu Lys Gly Thr Gln His Ile Met Gln Ala Gly Gln Thr 35 40 45

Leu His Leu Gln Cys Arg Gly Glu Ala Ala His Lys Trp Ser Leu Pro 50 55 60

Glu Met Val Ser Lys Glu Ser Glu Arg Leu Ser Ile Thr Lys Ser Ala 70 75 80

Cys Gly Arg Asn Gly Lys Gln Phe Cys Ser Thr Leu Thr Leu Asn Thr 85 90 95

Ala Gln Ala Asn His Thr Gly Phe Tyr Ser Cys Lys Tyr Leu Ala Val 100 105 110

Pro Thr Ser Lys Lys Lys Glu Thr Glu Ser Ala Ile Tyr Ile Phe Ile 115 120 125

Ser Asp Thr Gly Arg Pro Phe Val Glu Met Tyr Ser Glu Ile Pro Glu 130 135 140

Ile Ile His Met Thr Glu Gly Arg Glu Leu Val Ile Pro Cys Arg Val
145 150 155 160

Thr Ser Pro Asn Ile Thr Val Thr Leu Lys Lys Phe Pro Leu Asp Thr 165 170 175

Leu Ile Pro Asp Gly Lys Arg Ile Ile Trp Asp Ser Arg Lys Gly Phe 180 185 190

Ile Ile Ser Asn Ala Thr Tyr Lys Glu Ile Gly Leu Leu Thr Cys Glu

195 200 205

Ala	Thr 210	Val	Asn	Gly	His	Leu 215	Tyr	Lys	Thr	Asn	Tyr 220	Leu	Thr	His	Arg
Gln 225	Thr	Asn	Thr	Ile	Ile 230	Asp	Val	Gln	Ile	Ser 235	Thr	Pro	Arg	Pro	Val 240
Lys	Leu	Leu	Arg	Gly 245	His	Thr	Leu	Val	Leu 250	Asn	Cys	Thr	Ala	Thr 255	Thr
Pro	Leu	Asn	Thr 260	Arg	Val	Gln	Met	Thr 265	Trp	Ser	Tyr	Pro	Asp 270	Glu	Lys
Asn	Lys	Asn 275	Ala	Ser	Val	Arg	Arg 280	Arg	Ile	Asp	Gln	Ser 285	Asn	Ser	His
Ala	Asn 290	Ile	Phe	Tyr	Ser	Val 295	Leu	Thr	Ile	Asp	Lys 300	Met	Gln	Asn	Lys
Asp 305	Lys	Gly	Leu	Tyr	Thr 310	Cys	Arg	Val	Arg	Ser 315	Gly	Pro	Ser	Phe	Lys 320
Ser	Val	Asn	Thr	Ser 325	Val	His	Ile	Tyr	Asp 330	Lys	Ala	Gly	Pro	Gly 335	Glu
Pro	Lys	Ser	Cys 340	Asp	Lys	Thr	His	Thr 345	Cys	Pro	Pro	Cys	Pro 350	Ala	Pro
Glu	Leu	Leu 355	Gly	Gly	Pro	Ser	Val 360	Phe	Leu	Phe	Pro	Pro 365	Lys	Pro	Lys
Asp	Thr 370	Leu	Met		Ser			Pro	Glu	Val	Thr 380		Val	Val	Val
Asp 385	Val	Ser	His	Glu	Asp 390	Pro	Glu	Val	Lys	Phe 395	Asn	Trp	Tyr	Val	Asp 400
Gly	Val	Glu	Val	His 405	Asn	Ala	Lys	Thr	Lys 410	Pro	Arg	Glu	Glu	Gln 415	Tyr
Asn	Ser	Thr	Tyr	Arg	Val	Val	Ser	Val	Leu	Thr	Val	Leu	His	Gln	Asp

Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu 435 440 445

Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg 450 455 460

Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys 465 470 475 480

Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp 485 490 495

Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys 500 505 510

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser 515 520 525

Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser 530 540

Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser 545 550 555 560

Leu Ser Leu Ser Pro Gly Lys 565

<210> 21

<211> 1453

<212> DNA

<213> Artificial

<220>

<223> Flt 1 Receptor

<220>

<221> CDS

<222> (69)..(1442)

<400> 21 aagcttgggc tgcagg	tcga tcgactcta	g aggatcgatc	cccgggcgag ctcg	aattcg 60
caaccacc atg gtc Met Val 1	agc tac tgg gå Ser Tyr Trp As 5	c acc ggg gtc p Thr Gly Val	ctg ctg tgc gcg Leu Leu Cys Ala 10	g ctg 110 a Leu
ctc agc tgt ctg c Leu Ser Cys Leu L 15	tt ctc aca gga eu Leu Thr Gly 20	tct agt tcc (Ser Ser Ser (25	gga ggt aga cct Gly Gly Arg Pro	ttc 158 Phe 30
gta gag atg tac a Val Glu Met Tyr S 3	gt gaa atc ccc er Glu Ile Pro 5	gaa att ata Glu Ile Ile 1 40	cac atg act gaa His Met Thr Glu 45	gga 206 Gly
agg gag ctc gtc a Arg Glu Leu Val I 50				
act tta aaa aag t Thr Leu Lys Lys P 65				
ata atc tgg gac a Ile Ile Trp Asp S 80	gt aga aag ggc er Arg Lys Gly 85	Phe Ile Ile	tca aat gca acg Ser Asn Ala Thr 90	tac 350 Tyr
aaa gaa ata ggg c Lys Glu Ile Gly L 95				
tat aag aca aac t Tyr Lys Thr Asn T 1	at ctc aca cat yr Leu Thr His 15	cga caa acc Arg Gln Thr 2 120	aat aca atc ata Asn Thr Ile Ile 125	gat 446 Asp
gtg gtt ctg agt c Val Val Leu Ser P 130				
ctt gtc tta aat t Leu Val Leu Asn C 145		Thr Glu Leu		_
ttc aac tgg gaa t Phe Asn Trp Glu T 160		Lys His Gln		
aac cga gac cta a Asn Arg Asp Leu L 175				
agc acc tta act a Ser Thr Leu Thr I 1				
acc tgt gca gca t Thr Cys Ala Ala S 210				

								ggc Gly								782
								Gl ^A aaa								830
								atg Met								878
								cac His								926
aac Asn	tgg Trp	tac Tyr	gtg Val 290	gac Asp	ggc Gly	gtg Val	gag Glu	gtg Val 295	cat His	aat Asn	gcc Ala	aag Lys	aca Thr 300	aag Lys	ccg Pro	974
								tac Tyr								1022
_	_		_	_		_		ggc Gly	_			_	_	_	_	1070
								atc Ile								1118
								gtg Val								1166
								agc Ser 375								1214
								gag Glu								1262
								ccc Pro								1310
				-	-			gtg Val	-							1358
								atg Met								1406
	_	_	_	-			_	tct Ser				tgaç	gegge	ccg c	:	1453

450 455

<210> 22

<211> 458

<212> PRT

<213> Artificial

<400> 22

Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser 1 5 10 15

Cys Leu Leu Thr Gly Ser Ser Ser Gly Gly Arg Pro Phe Val Glu 20 25 30

Met Tyr Ser Glu Ile Pro Glu Ile Ile His Met Thr Glu Gly Arg Glu 35 40 45

Leu Val Ile Pro Cys Arg Val Thr Ser Pro Asn Ile Thr Val Thr Leu 50 60

Lys Lys Phe Pro Leu Asp Thr Leu Ile Pro Asp Gly Lys Arg Ile Ile 65 70 75 80

Trp Asp Ser Arg Lys Gly Phe Ile Ile Ser Asn Ala Thr Tyr Lys Glu 85 90 95

Ile Gly Leu Leu Thr Cys Glu Ala Thr Val Asn Gly His Leu Tyr Lys 100 105 110

Thr Asn Tyr Leu Thr His Arg Gln Thr Asn Thr Ile Ile Asp Val Val 115 120 125

Leu Ser Pro Ser His Gly Ile Glu Leu Ser Val Gly Glu Lys Leu Val 130 135 140

Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val Gly Ile Asp Phe Asn 145 150 155 160

Trp Glu Tyr Pro Ser Ser Lys His Gln His Lys Lys Leu Val Asn Arg 165 170 175

Asp	Leu	Lys	Thr 180	Gln	Ser	Gly	Ser	Glu 185	Met	Lys	Lys	Phe	Leu 190	Ser	Thr
Leu	Thr	Ile 195	Asp	Gly	Val	Thr	Arg 200	Ser	Asp	Gln	Gly	Leu 205	Tyr	Thr	Cys
Ala	Ala 210	Ser	Ser	Gly	Leu	Met 215	Thr	Lys	Lys	Asn	Ser 220	Thr	Phe	Val	Arg
Val 225	His	Glu	Lys	Gly	Pro 230	Gly	Asp	Lys	Thr	His 235	Thr	Cys	Pro	Pro	Cys 240
Pro	Ala	Pro	Glu	Leu 245	Leu	Gly	Gly	Pro	Ser 250	Val	Phe	Leu	Phe	Pro 255	Pro
Lys	Pro	Lys	Asp 260	Thr	Leu	Met	Ile	Ser 265	Arg	Thr	Pro	Glu	Val 270	Thr	Cys
Val	Val	Val 275	Asp	Val	Ser	His	Glu 280	Asp	Pro	Glu	Val	Lys 285	Phe	Asn	Trp
Tyr	Val 290	Asp	Gly	Val	Glu	Val 295	His	Asn	Ala	Lys	Thr 300	Lys	Pro	Arg	Glu
Glu 305	Gln	Tyr	Asn	Ser	Thr 310	Tyr	Arg	Val	Val	Ser 315	Val	Leu	Thr	Val	Leu 320
His	Gln	Asp	Trp	Leu 325	Asn	Gly	Lys	Glu	Tyr 330	Lys	Cys	Lys	Val	Ser 335	Asn
Lys	Ala		Pro 340				Glu			Ile	Ser		Ala 350		Gly
Gln	Pro	Arg 355	Glu	Pro	Gln	Val	Tyr 360	Thr	Leu	Pro	Pro	Ser 365	Arg	Asp	Glu
Leu	Thr 370	Lys	Asn	Gln	Val	Ser 375	Leu	Thr	Cys	Leu	Val 380	Lys	Gly	Phe	Tyr
Pro 385	Ser	Asp	Ile	Ala	Val 390	Glu	Trp	Glu	Ser	Asn 395	Gly	Gln	Pro	Glu	Asn 400
Asn	Tyr	Lys	Thr	Thr 405	Pro	Pro	Val	Leu	Asp 410	Ser	Asp	Gly	Ser	Phe 415	Phe

Leu	Tyr	Ser	Lys 420	Leu	Thr	Val	Asp	Lys 425	Ser	Arg	Trp	Gln	Gln 430	Gly	Asn	
Val	Phe	Ser 435	Cys	Ser	Val	Met	His 440	Glu	Ala	Leu	His	Asn 445	His	Tyr	Thr	
Gln	Lys 450	Ser	Leu	Ser	Leu	Ser 455	Pro	Gly	Lys							
<210	> 2	23														
<211	> :	L444														
<212	> I	ANC														
<213	> 1	Artii	icia	al												
<220	>															
<223	> I	Plt 1	Rec	cepto	or											
<220	>															
<221	> (CDS														
<222	>	(69).	. (14	136)												
<400		23														_,
_		_			-										aattcg	60
caac	caco														g ctg a Leu	110
ctc Leu 15																158
gta (Val																206
agg Arg																254
act Thr			_				-		-							302

					aga Arg											350
					ctg Leu 100											398
tat Tyr	aag Lys	aca Thr	aac Asn	tat Tyr 115	ctc Leu	aca Thr	cat His	cga Arg	caa Gln 120	acc Thr	aat Asn	aca Thr	atc Ile	ata Ile 125	gat Asp	446
					agg Arg											494
					acc Thr											542
					cca Pro											590
					caa Gln 180											638
					agc Ser											686
_					cag Gln	_										734
					ggc Gly											782
					Gly 999											830
					atg Met 260											878
					cac His											926
					gtg Val											974
cag Gln	tac Tyr	aac Asn	agc Ser	acg Thr	tac Tyr	cgt Arg	gtg Val	gtc Val	agc Ser	gtc Val	ctc Leu	acc Thr	gtc Val	ctg Leu	cac His	1022

305 310 315

						aag Lys 325										1070
						gag Glu										1118
ccc Pro	cga Arg	gaa Glu	cca Pro	cag Gln 355	gtg Val	tac Tyr	acc Thr	ctg Leu	ccc Pro 360	cca Pro	tcc Ser	cgg Arg	gat Asp	gag Glu 365	ctg Leu	1166
						ctg Leu										1214
_	_		_			tgg Trp		_			_	_				1262
						gtg Val 405										1310
						gac Asp										1358
		_			_	cat His		_								1406
_	_			_		ccg Pro			tga	gcgg	geege	2				1444

<210> 24

<211> 455

<212> PRT

<213> Artificial

<400> 24

Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser 1 5 10 15

Cys Leu Leu Thr Gly Ser Ser Ser Gly Gly Arg Pro Phe Val Glu 20 25 30

40 Leu Val Ile Pro Cys Arg Val Thr Ser Pro Asn Ile Thr Val Thr Leu Lys Lys Phe Pro Leu Asp Thr Leu Ile Pro Asp Gly Lys Arg Ile Ile 70 Trp Asp Ser Arg Lys Gly Phe Ile Ile Ser Asn Ala Thr Tyr Lys Glu 85 Ile Gly Leu Leu Thr Cys Glu Ala Thr Val Asn Gly His Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg Gln Thr Asn Thr Ile Ile Asp Ile Gln 115 120 Leu Leu Pro Arg Lys Ser Leu Glu Leu Leu Val Gly Glu Lys Leu Val 135 140 130 Leu Asn Cys Thr Val Trp Ala Glu Phe Asn Ser Gly Val Thr Phe Asp 160 145 150 155 Trp Asp Tyr Pro Gly Lys Gln Ala Glu Arg Gly Lys Trp Val Pro Glu 170 175 Arg Arg Ser Gln Gln Thr His Thr Glu Leu Ser Ser Ile Leu Thr Ile His Asn Val Ser Gln His Asp Leu Gly Ser Tyr Val Cys Lys Ala Asn 200 Asn Gly Ile Gln Arg Phe Arg Glu Ser Thr Glu Val Ile Val His Glu 215 Asn Gly Pro Gly Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro 230 Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys 245 250

Met Tyr Ser Glu Ile Pro Glu Ile Ile His Met Thr Glu Gly Arg Glu

Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val

265

260

Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp 275 280 285

Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr 290 295 300

Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp 305 310 315 320

Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu 325 330 335

Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg 340 345 350

Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys 355 360 365

Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp 370 375 380

Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys 385 390 395 400

Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser 405 410 415

Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser 420 425 430

Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser 435 440 445

Leu Ser Leu Ser Pro Gly Lys 450 455

<210> 25

<211> 1377

<212> DNA

<213> Artificial

<220>	
<223> Flt 1 Receptor	
<220>	
<221> CDS	
<222> (1)(1377)	
<400> 25 atg gtc agc tac tgg gac acc ggg gtc ctg ctg tgc gcg ctg ctc agc	48
Met Val Ser Tyr Trp Asp Thr Gly Val Leu Leu Cys Ala Leu Leu Ser 1 10 15	
tgt ctg ctt ctc aca gga tct agt tcc gga agt gat acc ggt aga cct Cys Leu Leu Thr Gly Ser Ser Ser Gly Ser Asp Thr Gly Arg Pro 20 25 30	96
ttc gta gag atg tac agt gaa atc ccc gaa att ata cac atg act gaa Phe Val Glu Met Tyr Ser Glu Ile Pro Glu Ile Ile His Met Thr Glu 35 40 45	144
gga agg gag ctc gtc att ccc tgc cgg gtt acg tca cct aac atc act Gly Arg Glu Leu Val Ile Pro Cys Arg Val Thr Ser Pro Asn Ile Thr 50 55 60	192
gtt act tta aaa aag ttt cca ctt gac act ttg atc cct gat gga aaa Val Thr Leu Lys Lys Phe Pro Leu Asp Thr Leu Ile Pro Asp Gly Lys 65 70 75 80	240
cgc ata atc tgg gac agt aga aag ggc ttc atc ata tca aat gca acg Arg Ile Ile Trp Asp Ser Arg Lys Gly Phe Ile Ile Ser Asn Ala Thr 85 90 95	288
tac aaa gaa ata ggg ctt ctg acc tgt gaa gca aca gtc aat ggg cat Tyr Lys Glu Ile Gly Leu Leu Thr Cys Glu Ala Thr Val Asn Gly His 100 105 110	336
ttg tat aag aca aac tat ctc aca cat cga caa acc aat aca atc ata Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg Gln Thr Asn Thr Ile Ile 115 120 125	384
gat gtg gtt ctg agt ccg tct cat gga att gaa cta tct gtt gga gaa Asp Val Val Leu Ser Pro Ser His Gly Ile Glu Leu Ser Val Gly Glu 130 135 140	432
aag ctt gtc tta aat tgt aca gca aga act gaa cta aat gtg ggg att Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val Gly Ile 145 150 155 160	480
gac ttc aac tgg gaa tac cct tct tcg aag cat cag cat aag aaa ctt Asp Phe Asn Trp Glu Tyr Pro Ser Ser Lys His Gln His Lys Lys Leu 165 170 175	528

					aaa Lys										576
_	_				ata Ile	_									624
					tcc Ser										672
					gaa Glu 230										720
					ctg Leu										768
		_	_		ctc Leu	_						-		_	816
					agc Ser										864
					gag Glu										912
	_			_	acg Thr 310		_	 _	_	-			_	_	960
	_	_		_	aat Asn		_		_	_	_	_			1008
	_			_	ccc Pro							_			1056
_		_	_		cag Gln			_					_		1104
_		_		_	gtc Val	_				_					1152
	_	_		_	gtg Val 390			 _			_		-		1200

Asn Tyr Lys	acc acg Thr Thr 405	cct ccc Pro Pro	gtg ctg Val Leu	gac tcc Asp Ser 410	gac ggc Asp Gly	tcc Ser	ttc Phe 415	ttc Phe	1248
ctc tac agc				Ser Arg					1296
gtc ttc tca Val Phe Ser 435						His			1344
cag aag agc Gln Lys Ser : 450									1377
<210> 26									
<211> 458									
<212> PRT									
<213> Artif	icial								
<400> 26									
<400> 26 Met Val Ser 1	Tyr Trp 5	Asp Thr	Gly Val	Leu Leu 10	Cys Ala	Leu	Leu 15	Ser	
Met Val Ser 1	5			10			15		
Met Val Ser 1	5 Leu Thr 20	Gly Ser	Ser Ser 25	10 Gly Ser	Asp Thr	Gly 30	15 Arg	Pro	
Met Val Ser 1 Cys Leu Leu 1	5 Leu Thr 20 Met Tyr	Gly Ser Ser Glu	Ser Ser 25 Ile Pro 40	Gly Ser	Asp Thr	Gly 30	15 Arg	Pro Glu	
Met Val Ser 1 Cys Leu Leu 1 Phe Val Glu 1 35	5 Leu Thr 20 Met Tyr Leu Val	Gly Ser Ser Glu Ile Pro	Ser Ser 25 Ile Pro 40 Cys Arg	Gly Ser	Asp Thr Ile His 45 Ser Pro 60	Gly 30 Met	15 Arg Thr	Pro Glu Thr	

Tyr Lys Glu Ile Gly Leu Leu Thr Cys Glu Ala Thr Val Asn Gly His
100 105 110

Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg Gln Thr Asn Thr Ile Ile Asp Val Val Leu Ser Pro Ser His Gly Ile Glu Leu Ser Val Gly Glu Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val Gly Ile Asp Phe Asn Trp Glu Tyr Pro Ser Ser Lys His Gln His Lys Lys Leu Val Asn Arg Asp Leu Lys Thr Gln Ser Gly Ser Glu Met Lys Lys Phe Leu Ser Thr Leu Thr Ile Asp Gly Val Thr Arg Ser Asp Gln Gly Leu Tyr Thr Cys Ala Ala Ser Ser Gly Leu Met Thr Lys Lys Asn Ser Thr Phe Val Arg Val His Glu Lys Asp Lys Thr His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu 355 360 365

Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr $370 \hspace{1cm} 375 \hspace{1cm} 380$

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn 385 390 395 400

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe 405 410 415

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn 420 425 430

Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr 435 440 445

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys 450 455

<210> 27

<211> 431

<212> PRT

<213> Artificial

<220>

<223> Peptide

<400> 27

Gly Arg Pro Phe Val Glu Met Tyr Ser Glu Ile Pro Glu Ile Ile His 1 10 15

Met Thr Glu Gly Arg Glu Leu Val Ile Pro Cys Arg Val Thr Ser Pro 20 25 30

Asn Ile Thr Val Thr Leu Lys Lys Phe Pro Leu Asp Thr Leu Ile Pro 35 40 45

Asp Gly Lys Arg Ile Ile Trp Asp Ser Arg Lys Gly Phe Ile Ile Ser 50 60

Asn Ala Thr Tyr Lys Glu Ile Gly Leu Leu Thr Cys Glu Ala Thr Val Asn Gly His Leu Tyr Lys Thr Asn Tyr Leu Thr His Arg Gln Thr Asn Thr Ile Ile Asp Val Val Leu Ser Pro Ser His Gly Ile Glu Leu Ser Val Gly Glu Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn 120 Val Gly Ile Asp Phe Asn Trp Glu Tyr Pro Ser Ser Lys His Gln His 135 Lys Lys Leu Val Asn Arg Asp Leu Lys Thr Gln Ser Gly Ser Glu Met 155 Lys Lys Phe Leu Ser Thr Leu Thr Ile Asp Gly Val Thr Arg Ser Asp Gln Gly Leu Tyr Thr Cys Ala Ala Ser Ser Gly Leu Met Thr Lys Lys 185 Asn Ser Thr Phe Val Arg Val His Glu Lys Gly Pro Gly Asp Lys Thr 195 His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser 215 Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg 235 Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr 295 Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu 330 Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Thr Cys Leu 345 340 Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn

360

Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser

370 375 380

Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg 385 390 395 400

Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu 405 410 415

His Asn His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys 420 425 430